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L8	L7 same L1	19	L8
L7	L6 near15 (efficien\$ or efficacy)	2100	L7
L6	(restriction near5 (digest or site or endonuclease or enzyme))	45576	L6
L5	(plasmid) near5 (multiple near5 polylinker)	16	L5
L4	(plasmid) near5 (two near2 polylinker)	3	L4
L3	(plasmid) near5 (two near5 polylinker)	30	L3
L2	(plasmid or vector) near5 ((two or multiple) near5 (polylinker or (cloning site)))	2316	L2
L1	(plasmid or vector) same ((two or multiple) near5 (polylinker or (cloning site)))	4865	L1

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<i>DB=USPT,PGPB,JPAB,EPAB,DWPI,TDBD; PLUR=YES; OP=ADJ</i>			
L2	plasmid same (two polylinker\$)	3	L2
L1	plasmid and (two polylinker\$)	8	L1

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Set	Items	Description
S1	81	(PLASMID?) (S) (POLYLINKER(3N) (MULTIPLE OR TWO))
S2	43	S1(S) (RESTRICTION(3N) (DIGEST? OR SITE? OR ENDONUCLEASE? OR ENZYME?))
S3	18	RD S2 (unique items)
S4	3	S3 (S) (EFFICIEN? OR EFFICACY)
S5	0	PLASMID(10N) (TWO(W) POLYLINKER?)
S6	1	PLASMID(10N) (MULTIPLE(W) POLYLINKER?)
S7	0	PLASMID(10N) (THREE(W) POLYLINKER?)
S8	0	PLASMID(S) (THREE(W) POLYLINKER?)
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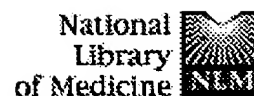
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
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Molecular cloning, phylogenetic analysis and three-dimensional modeling of Cu,Zn superoxide dismutase (CnSOD1) from three varieties of *Cryptococcus neoformans*.
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PMID: 11368899 [PubMed - indexed for MEDLINE]
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
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
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
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